Figure 1: Sequence listing of the OAS1 gene.

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1590 bp
                                                mRNA
                                                         linear
                                                                  PRI
            NM_016816
LOCUS
22-FEB-2001
DEFINITION Homo sapiens 2',5'-oligoadenylate synthetase 1 (40-46 kD)
(OAS1),
            transcript variant E18, mRNA.
            NM 016816
ACCESSION
            NM_016816.1 GI:8051620
VERSION
KEYWORDS
SOURCE
            human.
            Homo sapiens
  ORGANISM
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Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
Homo.
REFERENCE
            1 (bases 1 to 1590)
            Merlin, G., Chebath, J., Benech, P., Metz, R. and Revel, M.
  AUTHORS
            Molecular cloning and sequence of partial cDNA for
  TITLE
            interferon-induced (2'-5')oligo(A) synthetase mRNA from
human cells
            Proc. Natl. Acad. Sci. U.S.A. 80 (16), 4904-4908 (1983)
  JOURNAL
            83273721
  MEDLINE
            6348777
   PUBMED
            2 (bases 1 to 1590)
REFERENCE
            Saunders, M.E., Gewert, D.R., Tugwell, M.E., McMahon, M. and
  AUTHORS
            Williams, B.R.
            Human 2-5A synthetase: characterization of a novel cDNA
  TITLE
and
            corresponding gene structure
            EMBO J. 4 (7), 1761-1768 (1985)
  JOURNAL
            85284966
  MEDLINE
            2411547
   PUBMED
            3 (bases 1 to 1590)
REFERENCE
            Benech, P., Mory, Y., Revel, M. and Chebath, J.
  AUTHORS
            Structure of two forms of the interferon-induced (2'-5')
  TITLE
oligo A
            synthetase of human cells based on cDNAs and gene
sequences
            EMBO J. 4 (9), 2249-2256 (1985)
  JOURNAL
            86081732
  MEDLINE
   PUBMED
            2416561
REFERENCE
            4 (bases 1 to 1590)
            Wathelet, M., Moutschen, S., Cravador, A., DeWit, L.,
  AUTHORS
Defilippi, P.,
            Huez, G. and Content, J.
             Full-length sequence and expression of the 42 kDa 2-5A
  TITLE
synthetase
             induced by human interferon
             FEBS Lett. 196 (1), 113-120 (1986)
  JOURNAL
             86108911
  MEDLINE
             3753689
   PUBMED
             5 (bases 1 to 1590)
REFERENCE
             Shiojiri, S., Fukunaga, R., Ichii, Y. and Sokawa, Y.
  AUTHORS
             Structure and expression of a cloned cDNA for human
  TITLE
             (2'-5')oligoadenylate synthetase
             J. Biochem. 99 (5), 1455-1464 (1986)
  JOURNAL
             86223945
  MEDLINE
   PUBMED
             3754863
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PCT/GB03/01625

Figure 1:Page 2

(bases 1 to 1590) REFERENCE Williams, B.R., Saunders, M.E. and Willard, H.F. AUTHORS Interferon-regulated human 2-5A synthetase gene maps to TITLE chromosome 12 Somat. Cell Mol. Genet. 12 (4), 403-408 (1986) JOURNAL 86289724 MEDLINE 2426799 PUBMED (bases 1 to 1590) REFERENCE Benech, P., Vigneron, M., Peretz, D., Revel, M. and AUTHORS Chebath, J. Interferon-responsive regulatory elements in the promoter TITLE of the human 2',5'-oligo(A) synthetase gene Mol. Cell. Biol. 7 (12), 4498-4504 (1987) JOURNAL 88142842 MEDLINE 2830497 PUBMED (bases 1 to 1590) REFERENCE Wathelet, M.G., Clauss, I.M., Nols, C.B., Content, J. and AUTHORS Huez, G.A. New inducers revealed by the promoter sequence analysis TITLE of two interferon-activated human genes Eur. J. Biochem. 169 (2), 313-321 (1987) JOURNAL 88082760 MEDLINE 3121313 PUBMED 9 (bases 1 to 1590) REFERENCE Rutherford, M.N., Hannigan, G.E. and Williams, B.R. AUTHORS Interferon-induced binding of nuclear factors to promoter TITLE elements of the 2-5A synthetase gene EMBO J. 7 (3), 751-759 (1988) JOURNAL MEDLINE 88283644 PUBMED 2456211 10 (bases 1 to 1590) REFERENCE Wathelet, M.G., Szpirer, J., Nols, C.B., Clauss, I.M., De AUTHORS Wit, L., Islam, M.Q., Levan, G., Horisberger, M.A., Content, J., Szpirer, C. and Huez, G.A. Cloning and chromosomal location of human genes inducible TITLE by type I interferon Somat. Cell Mol. Genet. 14 (5), 415-426 (1988) JOURNAL 89019578 MEDLINE 3175763 PUBMED 11 (bases 1 to 1590) REFERENCE Nechiporuk T, Nechiporuk A, Sahba S, Figueroa K, Shibata AUTHORS H, Chen XN, Korenberg JR, de Jong P and Pulst SM. A high-resolution PAC and BAC map of the SCA2 region TITLE Genomics 44 (3), 321-329 (1997) JOURNAL 97468145 MEDLINE 9325053 PUBMED 12 (bases 1 to 1590) REFERENCE

Figure 1:Page 3

AUTHORS Renault B, Hovnanian A, Bryce S, Chang JJ, Lau S, Sakuntabhai A,
Monk S, Carter S, Ross CJ, Pang J, Twells R, Chamberlain S, Monaco

AP, Strachan T and Kucherlapati R. A sequence-ready physical map of a region of 12q24.1 TITLE Genomics 45 (2), 271-278 (1997) JOURNAL 98008914 MEDLINE 9344649 PUBMED 13 (bases 1 to 1590) REFERENCE Hovnanian, A., Rebouillat, D., Mattei, M.G., Levy, E.R., AUTHORS Marie, I., Monaco, A.P. and Hovanessian, A.G. The human 2',5'-oligoadenylate synthetase locus is TITLE composed of three distinct genes clustered on chromosome 12q24.2 encoding the 100-, 69-, and 40-kDa forms Genomics 52 (3), 267-277 (1998) JOURNAL 99009311 MEDLINE 9790745 PUBMED REVIEWED REFSEQ: This record has been curated by NCBI COMMENT staff. The reference sequence was derived from X02875.1. Summary: This gene encodes an enzyme included in the 2', 5 ' oligoadenylate synthase family. This enzyme is induced bу interferons and catalyzes the 2', 5' oligomers of adenosine in order to bind and activate RNase L. This enzyme family plays a significant role in the inhibition of cellular protein synthesis and viral infection resistance. Alternative splicing of this gene produces two isoforms which share identical N-terminal sequence but diverge at exon 7. Transcript Variant: Transcript variant E18 represents the 1.8 kb product that diverges from the predominant transcript variant E16 in exon 7. This form contains an additional exon and is very hydrophilic as compared to transcript variant E16. COMPLETENESS: complete on the 3' end. **FEATURES** Location/Qualifiers 1..1590 source /organism="Homo sapiens" /db xref="taxon:9606" /chromosome="12" /map="12q24.1" 1..1590 gene /gene="OAS1"

/note="IFI-4; OIAS"

Figure 1:Page 4

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polyA_signal
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polyA_site
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124

Figure 1:Page 5

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|---------------------------------|------------|------------|------------|------------|------------|
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| | aaatctctgg | acaagttcat | tgaagactat | ctcttgccag | acacgtgttt |
| | atcgaccatg | ccattgacat | catctgtggg | ttcctgaagg | aaaggtgctt |
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| ggaagcctgt 361 gggcaacccc | caaagagaga | gagcactttc | cgtgaagttt | gaggtccagg | ctccacgctg |
| 421 gttcgatgtg | cgtgcgctca | gcttcgtact | gagttcgctc | cagctcgggg | agggggtgga |
| 481 cccccaaato | ctgcctgcct | ttgatgccct | gggtcagttg | actggcagct | ataaacctaa |
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| | ttcacagaac | tacagagaga | cttcctgaag | cagcgcccca | ccaagctcaa |
| 661 qccacctcag | cgcctagtca | agcactggta | ccaaaattgt | aagaagaagc | ttgggaagct |
| | tatgccctgg | agctcctgac | ggtctatgct | tgggagcgag | ggagcatgaa |
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| | cagctcacga | aacccaggcc | tgtgatcctg | gacccggcgg | accctacagg |
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| 1021 ggctgaaago | ccatgcttta | agaattggga | | | |
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| atccacccca | 1 | accctcattt | | | |
| tgggggaaac | 1 | | | | cagtgcatct |
| atccagagaa | 1 | | | | gggactcttg |
| tcctgactcc | 2 | | | | aacccaagtc |
| acttcattco | 3 | | | | ccacagcctc |
| gagaatgaaa | 3. | | | | atttagataa |
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Figure 2: Schematic of the OAS1 gene showing intron/exon structure and protein structure

Genomic structure

